PCT

RAW SEQUENCE LISTING   DATE: 09/09/2004
PATENT APPLICATION: US/10/506,455   TIME: 16:17:57

Input Set : **A:\X-15648.ST25.txt**
Output Set: **N:\CRF4\09092004\J506455.raw**

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 3 <110> APPLICANT: Beals, John
 4         Kuchibhotla, Uma
 6 <120> TITLE OF INVENTION: HETEROLOGOUS G-CSF FUSION PROTEINS
 8 <130> FILE REFERENCE: X-15648
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C--> 10 **<140> CURRENT APPLICATION NUMBER: US/10/506,455**
C--> 10 **<141> CURRENT FILING DATE: 2004-08-31**

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11 <151> PRIOR FILING DATE: 2003-02-21
13 <160> NUMBER OF SEQ ID NOS: 66
15 <170> SOFTWARE: PatentIn version 3.2
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18 <211> LENGTH: 174
19 <212> TYPE: PRT
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39 <223> OTHER INFORMATION: Xaa at position 38 is Thr, or any other amino acid exept Pro;
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53 <222> LOCATION: (58)..(58)
54 <223> OTHER INFORMATION: Xaa at position 58 is Trp or Asn;
56 <220> FEATURE:
57 <221> NAME/KEY: MISC_FEATURE
58 <222> LOCATION: (59)..(59)
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    68 <222> LOCATION: (61)..(61)
    69 <223> OTHER INFORMATION: Xaa at position 61 is Leu, or any other amino acid except
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    72 <221> NAME/KEY: MISC_FEATURE
    73 <222> LOCATION: (62)..(62)
    74 <223> OTHER INFORMATION: Xaa at position 62 is Ser or Thr;
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    78 <222> LOCATION: (63)..(63)
    79 <223> OTHER INFORMATION: Xaa at position 63 Ser or Asn;
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    83 <222> LOCATION: (64)..(64)
    84 <223> OTHER INFORMATION: Xaa at position 64 is Cys or any other amino acid except Pro;
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    94 <223> OTHER INFORMATION: Xaa at position 66 is Ser or Thr;
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    99 <223> OTHER INFORMATION: Xaa at position 67 is Gln or Asn;
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   104 <223> OTHER INFORMATION: Xaa at position 68 is Ala or any other amino acid except
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   114 <223> OTHER INFORMATION: Xaa at position 93 is Glu or Asn;
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   118 <222> LOCATION: (94)..(94)
   119 <223> OTHER INFORMATION: Xaa at position 94 is Gly or any other amino acid except
Pro;
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122 <221> NAME/KEY: MISC_FEATURE
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124 <223> OTHER INFORMATION: Xaa at position 95 is Ile, Asn, Ser, or Thr;
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134 <223> OTHER INFORMATION: Xaa at position 133 is Thr or Asn;
136 <220> FEATURE:
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139 <223> OTHER INFORMATION: Xaa at position 134 is Gln or any other amino acid except
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154 <223> OTHER INFORMATION: Xaa at position 142 is Ser or any other amino acid except
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159 <223> OTHER INFORMATION: Xaa at position 143 is Ala, Ser, or Thr.
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168              20                  25                  30
171 Glu Lys Leu Cys Xaa Xaa Xaa Lys Leu Cys His Pro Glu Glu Leu Val
172          35                  40                  45
175 Leu Leu Gly His Ser Leu Gly Ile Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
176      50                  55                  60
179 Xaa Xaa Xaa Xaa Xaa Gln Leu Ala Gly Cys Leu Ser Gln Leu His Ser
180 65                  70                  75                  80
183 Gly Leu Phe Leu Tyr Gln Gly Leu Leu Gln Ala Leu Xaa Xaa Xaa Ser
184              85                  90                  95
187 Xaa Glu Leu Gly Pro Thr Leu Asp Thr Leu Gln Leu Asp Val Ala Asp
188              100                 105                 110
191 Phe Ala Thr Thr Ile Trp Gln Gln Met Glu Glu Leu Gly Met Ala Pro
192          115                 120                 125
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195 Ala Leu Gln Pro Xaa Xaa Xaa Ala Met Pro Ala Phe Xaa Xaa Xaa Phe
196         130             135             140
199 Gln Arg Arg Ala Gly Gly Val Leu Val Ala Ser His Leu Gln Ser Phe
200 145             150             155             160
203 Leu Glu Val Ser Tyr Arg Val Leu Arg His Leu Ala Gln Pro
204             165             170
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208 <211> LENGTH: 1044
209 <212> TYPE: DNA
210 <213> ORGANISM: Artificial Sequence
212 <220> FEATURE:
213 <223> OTHER INFORMATION: synthetic construct
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220 cagggcgatg gcgcagcgct ccagcggaat ctcgttcact ccttctaggt cccgctaccg    180
222 cgtcgcgagg tcgagaagct gtgtgccacc tacaagctgt gccaccccga ggagctggtg    240
224 ctcttcgaca cacggtggat gttcgacacg gtggggctcc tcgaccacct gctcggacac    300
226 tctctgggca tcccctgggc tccctgagc agctgcgacg agcctgtgac agacccgtag    360
228 gggacccgag gggactcgtc gacgcccagc caggccctgc agctggcagg ctgcttgagc    420
230 caactccata gcgggtcggt ccgggacgtc gaccgtccga cgaactcggt tgaggtatcg    480
232 ggccttttcc tctaccaggg gctcctgcag gccctggaag ggatctcccc ggaaaaggag    540
234 atggtccccg aggacgtccg ggaccttccc tagaggcccg agttgggtcc caccttggac    600
236 acactgcagc tggacgtcgc cgacgggctc aacccagggt ggaacctgtg tgacgtcgac    660
238 ctgcagcggc tgtttgccac caccatctgg cagcagatgg aagaactggg aatggcccct    720
240 aaacggtggt ggtagaccgt cgtctacctt cttgaccctt accggggagc cctgcagccc    780
242 aaccagaccg ccatgccggc cttcgcctct gctttccggg acgtcgggtt ggtctggcgg    840
244 tacggccgga agcggagacg aaagcagcgc cgggcaggag gggtcctggt tgcctcccat    900
246 ctgcagagct tcgtcgcggc ccgtcctccc caggaccaac ggagggtaga cgtctcgaag    960
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256 <213> ORGANISM: Artificial Sequence
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268 cgtcgcgagg tcgagaagct gtgtgccacc tacaagctgt gccaccccga ggagctggtg    240
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274 gggacccgag gggactcgtc gacgcccagc caggccctgc agctggcagg ctgcttgagc    420
276 caactccata gcgggtcggt ccgggacgtc gaccgtccga cgaactcggt tgaggtatcg    480
278 ggccttttcc tctaccaggg gctcctgcag gccctggaag ggatctcccc ggaaaaggag    540
280 atggtccccg aggacgtccg ggaccttccc tagaggcccg agttgggtcc caccttggac    600
282 acactgcagc tggacgtcgc cgacgggctc aacccagggt ggaacctgtg tgacgtcgac    660
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288 acccagggtg ccatgccggc cttcaactct accttccggg acgtcgggtg ggtcccacgg    840
290 tacggccgga agttgagatg gaagcagcgc cgggcaggag gggtcctggt tgcctcccat    900
292 ctgcagagct tcgtcgcggc ccgtcctccc caggaccaac ggagggtaga cgtctcgaag    960
294 ctggaggtgt cgtaccgcgt cttaaggcac cttgcccagc ccgacctcca cagcatggcg   1020
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302 <213> ORGANISM: Artificial Sequence
304 <220> FEATURE:
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314 cgtcgcgagg tcgagaagct gtgtaacacc accaagctgt gccaccccga ggagctggtg     240
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320 gggacccgag gggactcgtc gacgcccagc caggccctgc agctggcagg ctgcttgagc     420
322 caactccata gcgggtcggt ccgggacgtc gaccgtccga cgaactcggt tgaggtatcg     480
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326 atggtccccg aggacgtccg ggaccttccc tagaggcccg agttgggtcc caccttggac     600
328 acactgcagc tggacgtcgc cgacgggctc aacccagggt ggaacctgtg tgacgtcgac     660
330 ctgcagcggc tgtttgccac caccatctgg cagcagatgg aagaactggg aatggcccct     720
332 aaacggtggt ggtagaccgt cgtctacctt cttgacccttt accggggagc cctgcagccc     780
334 acccagggtg ccatgccggc cttcgcctct gctttccggg acgtcgggtg ggtcccacgg     840
336 tacggccgga agcggagacg aaagcagcgc cgggcaggag gggtcctggt tgcctcccat     900
338 ctgcagagct tcgtcgcggc ccgtcctccc caggaccaac ggagggtaga cgtctcgaag     960
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347 <212> TYPE: DNA
348 <213> ORGANISM: Artificial Sequence
350 <220> FEATURE:
351 <223> OTHER INFORMATION: synthetic construct
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358 cagggcgatg gcgcagcgct ccagcggaat ctcgttcact ccttctaggt cccgctaccg     180
360 cgtcgcgagg tcgagaagct gtgtgccacc tacaagctgt gccaccccga ggagctggtg     240
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364 tctctgggca tcccctgggc taacactagc agctgcgacg agcctgtgac agacccgtag     360
366 gggacccgat tggactcctc gacgcccagc caggccctgc agctggcagg ctgcttgagc     420
368 caactccata gcgggtcggt ccgggacgtc gaccgtccga cgaactcggt tgaggtatcg     480
370 ggcctttttcc tctaccaggg gctcctgcag gccctggaag ggatctcccc ggaaaaggag     540
372 atggtccccg aggacgtccg ggaccttccc tagaggcccg agttgggtcc caccttggac     600
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RAW SEQUENCE LISTING ERROR SUMMARY　　DATE: 09/09/2004
PATENT APPLICATION:　US/10/506,455　　TIME: 16:17:58

Input Set : **A:\X-15648.ST25.txt**
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<u>Please Note:</u>
**Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.**

Seq#:1; Xaa Pos. 17,37,38,39,57,58,59,60,61,62,63,64,65,66,67,68,69,93,94
Seq#:1; Xaa Pos. 95,97,133,134,135,141,142,143

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:167 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:16
M:341 Repeated in SeqNo=1